

ROBUST SCIENCE WORKSHOP
FEBRUARY 17, 2021

TOWARDS AN HPC AUTOMATED PIPELINE FOR CONNECTOMICS

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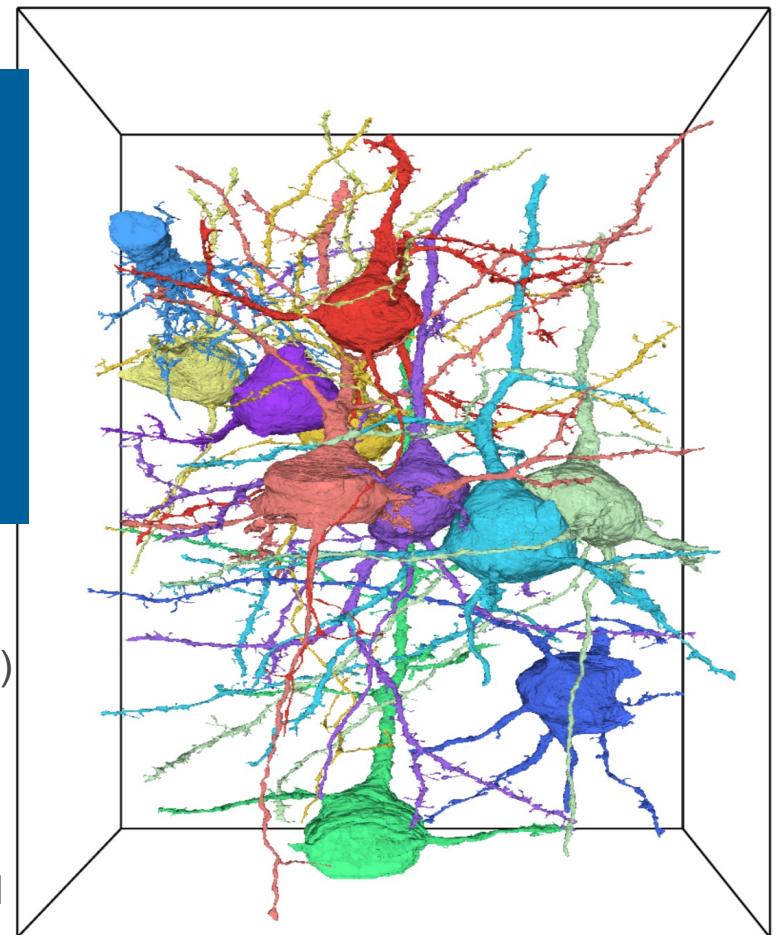
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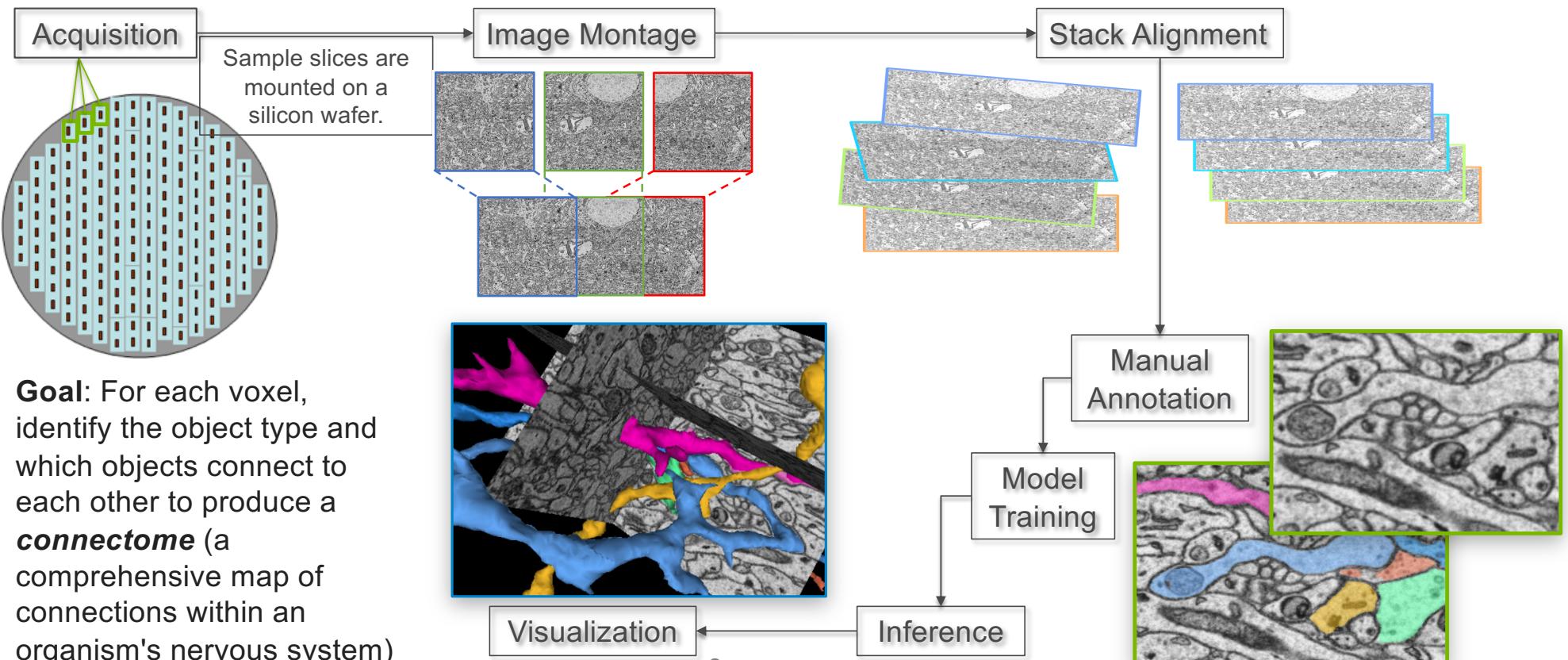
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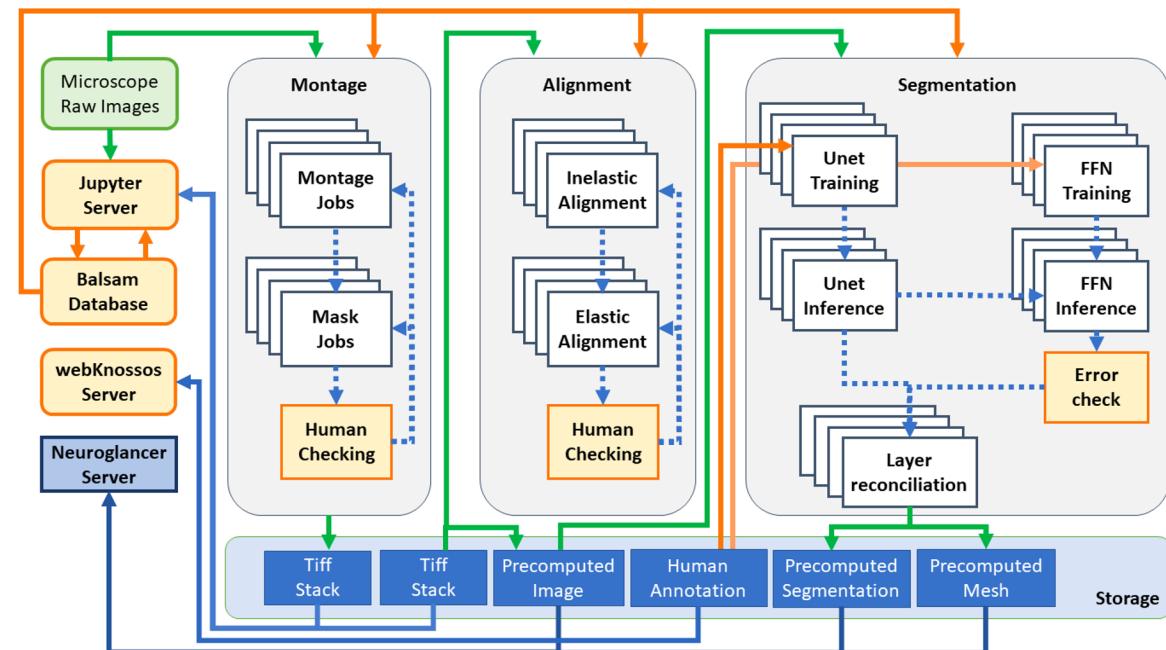
SERIAL ELECTRON MICROSCOPY ACQUISITION AND ANALYSIS STEPS FOR CONNECTOMICS



Scalable, trustworthy, and reproducible

Our experience: EM PIPELINE @ ARGONNE

- Optimizing codes for montage, alignment, and segmentation to run on Argonne supercomputers with natural concurrency (tens to thousands of compute nodes)
- Assembling codes into a pipeline
 - Respect (and exploit) concurrency of individual applications
 - **Reduce human-intensive aspects of large-scale computing**
 - Schedule jobs in a more optimal/efficient fashion
 - Support user interaction with data throughout pipeline (orange boxes)



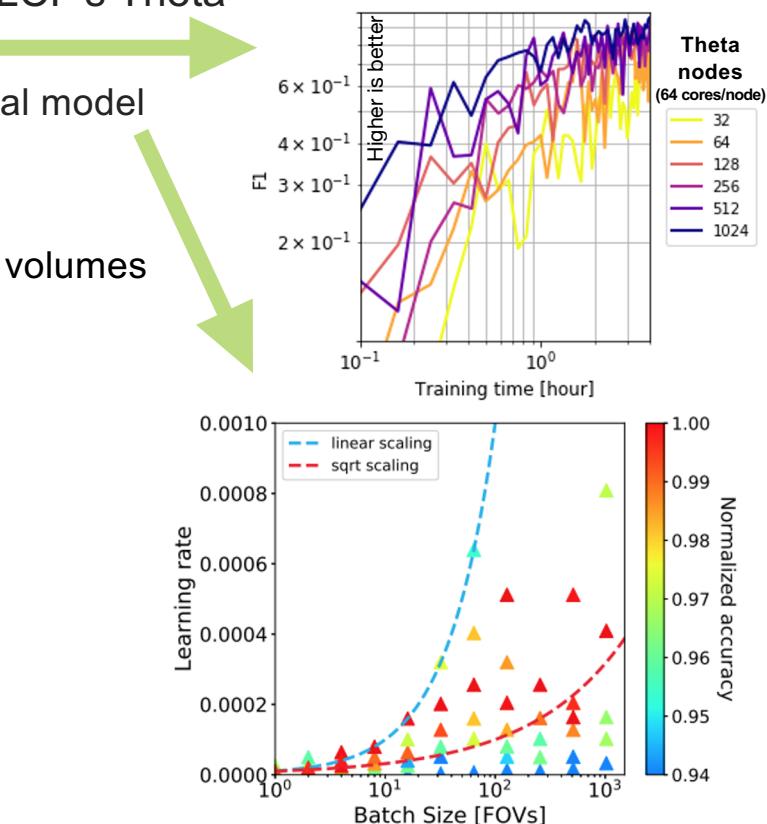
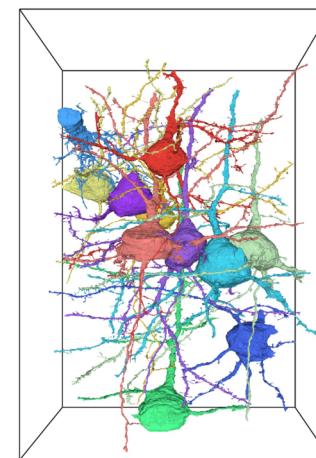
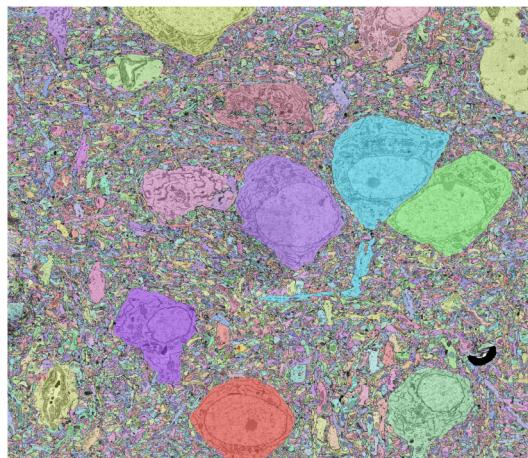
LARGE-SCALE FLOOD FILLING NETWORKS (FFN)

■ Scaled training of Google's FFN to thousands of nodes on ALCF's Theta

- Large-scale training achieves greater accuracy faster
- Faster training opens opportunity to make many runs to reach optimal model (hyperparameter optimization)

■ FFN Segmentation rate on Theta: ~350M voxels/nodehour

- Distributed inference on many nodes allows us to reconstruct larger volumes faster (weak scaling)



WORKSHOP QUESTION

Can publishing/dissemination/sharing standard practices and procedures contribute to achieving or encouraging scalability, trust, and reproducibility in the applications results?

- Neuroscience computational community shares procedures, practices, data
- Scalability being addressed by some
- Trust – from human evaluation – bottleneck!

Main challenges:

- Reducing human effort (proof reading)
 - Develop better metrics for intermediate steps

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